Attorney Docket No. 208859

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

In re Application of:

Roelvink et al.

Group Art Unit: Unknown

Application No. unassigned

Examiner: Unknown

Filing Date: February 9, 2001

Date: February 9, 2001

For: ADENOVIRAL CAPSID CONTAINING

CHIMERIC PROTEIN IX

SUBMISSION OF SEQUENCE LISTING

In accordance with the requirements of 37 CFR 1.821-1.825, a nucleotide/amino acid sequence listing is submitted as part of the new patent application identified above. A sequence listing in written form (paper copy), with pages numbered separately from the pages of the application, is enclosed. A sequence listing in a computer readable version (diskette) that is identical to the sequence listing in written form is also enclosed. The undersigned agent verifies that the paper copy of the sequence listing and the computer readable version of the sequence listing are identical.

Respectfully submitted,

By

John Rilyk, Jr., Reg No. 35,763

Attorney for the Applicant

LEYDIG, VOIT & MAYER, LTD. Two Prudential Plaza, Suite 4900

180 North Stetson

Chicago, Illinois 60601-6780 (US)

(312) 616-5600 Telephone

(312) 616-5700 Facsimile

بالملالين

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SEQUENCE LISTING

<110> Roelvink, Petrus W Kovesdi, Imre Wickham, Thomas J

<120> ADENOVIRAL CAPSID CONTAINING CHIMERIC PROTEIN IX

<130> 208859

<140> US

<141> 2001-02-09

<150> US 60/181,163

<151> 2000-02-09

<160> 13

<170> PatentIn Ver. 2.1

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Gln Asn Val Val Gly Ser Thr Val Asp Gly Arg Pro Val Ala Pro Ala 35 40 45

Asn Ser Ser Thr Leu Thr Tyr Ala Thr Ile Gly Pro Ser Pro Leu Asp
50 55 60

Thr Ala Ala Ala Ala Ala Ser Ala Ala Ala Ser Thr Ala Arg Ser 65 70 75 80

Met Ala Ala Asp Phe Ser Phe Tyr Asn His Leu Ala Ser Asn Ala Val 85 90 95

Thr Arg Thr Ala Val Arg Glu Asp Ile Leu Thr Val Met Leu Ala Lys
100 105 110

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Leu Leu Ala Thr Leu Asn Arg Leu Arg Thr Gly Leu Ala Ala Tyr Val

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Gln Ala Asn Leu Val Gly Gly Gln Val Asn Pro Phe Val 115 120 125

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Thr Arg Met Pro Pro Trp Ala Gly Val Arg Gln Asn Val Met Gly Ser 20 25 30

Ser Ile Asp Gly Arg Pro Val Leu Pro Ala Asn Ser Thr Thr Leu Thr 35 40 45

Tyr Glu Thr Val Ser Gly Thr Pro Leu Glu Thr Ala Ala Ser Ala Ala 50 55 60

Ala Ser Ala Ala Ala Ala Thr Ala Arg Gly Ile Val Thr Asp Phe Ala 65 70 75 80

Phe Leu Ser Pro Leu Ala Ser Ser Ala Ala Ser Arg Ser Ser Ala Arg
85 90 95

Asp Asp Lys Leu Thr Ala Leu Leu Ala Gln Leu Asp Ser Leu Thr Arg
100 105 110

Glu Leu Asn Val Val Ser Gln Gln Leu Leu Asp Leu Arg Gln Gln Val
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Ser Ala Leu Lys Ala Ser Ser Pro Pro Asn Ala Val 130 135 140

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Tyr Glu Thr Val Ser Gly Thr Pro Leu Glu Thr Ala Ala Ser Ala Ala 50 55 60

Ala Ser Ala Ala Ala Ala Thr Ala Arg Gly Ile Val Thr Asp Phe Ala 65 70 75 80

Phi

4

Phe Leu Ser Pro Leu Ala Ser Ser Ala Ala Ser Arg Ser Ser Ala Arg 85 90 95

Asp Asp Lys Leu Thr Ala Leu Leu Ala Gln Leu Asp Ser Leu Thr Arg 100 105 110

Glu Leu Asn Val Val Ser Gln Gln Leu Leu Asp Leu Arg Gln Gln Val 115 120 125

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20 25 30

Gln Asn Val Gly Ser Asn Val Asp Gly Arg Pro Val Ala Pro Ala 35 40 45

Asn Ser Thr Thr Leu Thr Tyr Ala Thr Ile Gly Ser Ser Val Asp Thr 50 55 60

Ala Ala Ala Ala Ala Ser Ala Ala Ser Thr Ala Arg Gly Met 65 70 75 80

Ala Ala Asp Phe Gly Leu Tyr Asn Gln Leu Ala Ala Ser Arg Leu Arg 85 90 95

Glu Glu Asp Ala Leu Ser Val Val Leu Thr Arg Leu Glu Glu Leu Ser

Gln Gln Leu Gln Asp Met Ser Ala Lys Met Ala Leu Leu Asn Pro Pro 115 120 125

Ala Asn Thr Ser

<210> 7

<211> 133

<212> PRT

<213> Adenovirus

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1 5 10 15

Phe Ser Pro Tyr Leu Thr Thr Arg Leu Pro Ala Trp Ala Gly Val Arg
20 25 30

t Miles

Gln Asn Val Met Gly Ser Asn Val Asp Gly Arg Pro Val Ala Pro Ala 35 40 45

Asn Ser Ala Thr Leu Thr Tyr Ala Thr Val Gly Ser Ser Val Asp Thr 50 55 60

Ala Ala Ala Ala Ala Ser Ala Ala Ser Thr Ala Arg Gly Met 65 70 75 80

Ala Ala Asp Phe Gly Leu Tyr Asn Gln Leu Ala Ala Ser Arg Ser Leu 85 90 95

Arg Glu Glu Asp Ala Leu Ser Val Val Leu Thr Arg Met Glu Glu Leu
100 105 110

Ser Gln Gln Leu Gln Asp Leu Phe Ala Lys Val Ala Leu Leu Asn Pro 115 120 125

Pro Ala Asn Ala Ser 130

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Asn Xaa Asp Gly Arg Pro Val Leu Pro Ala Asn Ser Xaa Thr Leu Thr 35 40 45

Tyr Glu Thr Val Gly Xaa Xaa Xaa Xaa Thr Ala Ala Ala Ala Ala 50 55 60

Ser Ala Ala Ala Xaa Thr Ala Arg Gly Xaa Xaa Asp Phe Xaa Xaa 65 70 75

Xaa Xaa Xaa Leu Ala Xaa Ser Xaa Xaa Xaa Arg Xaa Xaa Xaa Glu 85 90 95

Asp Xaa Leu Xaa Xaa Leu Leu Ala Xaa Leu Xaa Xaa Leu Xaa Xaa Xaa 100 105 110

Leu Xaa Xaa Xaa Ser Gln Xaa Xaa Leu Xaa Xaa Xaa Pro Xaa Asn 115 120 125

Xaa Val

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<211> 130 <212> PRT

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<212> PRT

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<400> 11

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Gly Ser Gly Ser Gly Ser Thr Arg Ser Thr Asn Ser Phe Asp
20 25 30

1111

Gly Ser Ile Val Ser Ser Tyr Leu Thr Thr Arg Met Pro Pro Trp Ala Gly Val Arg Gln Asn Val Met Gly Ser Ser Ile Asp Gly Arg Pro Val Leu Pro Ala Asn Ser Thr Thr Leu Thr Tyr Glu Thr Val Ser Gly Thr 70 Pro Leu Glu Thr Ala Ala Ser Ala Ala Ala Ser Ala Ala Ala Thr Ala Arg Gly Ile Val Thr Asp Phe Ala Phe Leu Ser Pro Leu Ala Ser 100 105 Ser Ala Ala Ser Arg Ser Ser Ala Arg Asp Asp Lys Leu Thr Ala Leu Leu Ala Gln Leu Asp Ser Leu Thr Arg Glu Leu Asn Val Val Ser Gln 130 135 Gln Leu Leu Asp Leu Arg Gln Gln Val Ser Ala Leu Lys Ala Ser Ser 145 150 155 Pro Pro Asn Ala Val 165 <210> 12 <211> 495 <212> DNA <213> Adenovirus <400> 12 atgagcacca actogtttga tggaagcatt gtgagctcat atttgacaac gcgcatgccc 60 ccatgggccg gggtgcgtca gaatgtgatg ggctccagca ttgatggtcg ccccgtcctg 120 cccgcaaact ctactacctt gacctacgag accgtgtctg gaacgccgtt ggagactgca 180 gcctccgccg ccgcttcagc cgctgcagcc accgcccgcg ggattgtgac tgactttgct 240 ttcctgagcc cgcttgcaag cagtgcagct tcccgttcat ccgcccgcga tgacaagttg 300 acggetettt tggcacaatt ggattetttg acccgggaac ttaatgtegt tteteageag 360 tctagtggtt ctggctcagg ctccggttca ggttcgggat cttaccccta cgacgtgccc 480 gactacgcct ctaga 495 <210> 13 <211> 165 <212> PRT <213> Adenovirus <400> 13 Met Ser Thr Asn Ser Phe Asp Gly Ser Ile Val Ser Ser Tyr Leu Thr

10

Thr Arg Met Pro Pro Trp Ala Gly Val Arg Gln Asn Val Met Gly Ser

25

20

.22.

- Ser Ile Asp Gly Arg Pro Val Leu Pro Ala Asn Ser Thr Thr Leu Thr 35
- Tyr Glu Thr Val Ser Gly Thr Pro Leu Glu Thr Ala Ala Ser Ala Ala 50 60
- Ala Ser Ala Ala Ala Ala Thr Ala Arg Gly Ile Val Thr Asp Phe Ala 65 70 75 80
- Phe Leu Ser Pro Leu Ala Ser Ser Ala Ala Ser Arg Ser Ser Ala Arg
- Asp Asp Lys Leu Thr Ala Leu Leu Ala Gln Leu Asp Ser Leu Thr Arg 100 105 110
- Glu Leu Asn Val Val Ser Gln Gln Leu Leu Asp Leu Arg Gln Gln Val
- Ser Ala Leu Lys Ala Ser Ser Pro Pro Asn Ala Val Ser Ser Gly Ser 130 135 140
- Gly Ser Gly Ser Gly Ser Gly Ser Tyr Pro Tyr Asp Val Pro 145 150 155 160

Asp Tyr Ala Ser Arg 165